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RAW SEQUENCE LISTING DATE: 04/17/2001 PATENT APPLICATION: US/09/691,763B TIME: 13:21:40

Input Set : A:\E03557003.txt

Output Set: N:\CRF3\04172001\1691763B.raw

egeaaceagg geagggeagg geagggeagg caeggettgg caaccetgeg caececacee

accaacccac accetgeggg ggaagggaga caatattace etcateccae tgcatgtggg

4 <110> APPLICANT: Vertino, Paula M. ENTERED See p. 5 6 <120> TITLE OF INVENTION: TMS1 Compositions and Methods of Use 9 <130> FILE REFERENCE: E0355/7003/ERG/MAT 11 <140> CURRENT APPLICATION NUMBER: US 09/691,763B 12 <141> CURRENT FILING DATE: 2000-10-18 14 <150> PRIOR APPLICATION NUMBER: US 60/159,975 15 <151> PRIOR FILING DATE: 1999-10-18 17 <160> NUMBER OF SEQ ID NOS: 27 19 <170> SOFTWARE: FastSEQ for Windows Version 3.0 21 <210> SEQ ID NO: 1 22 <211> LENGTH: 2821 23 <212> TYPE: DNA 24 <213> ORGANISM: Homo Sapiens 26 <400> SEQUENCE: 1 27 aagettttga atacaacatg ctgcaggcat cacagcctca ttcattcctt cactcagcaa 60 120 28 atctttactc agcacctaat gtgttccaga tacatttttt tttttcagat ggaatctagc totgtcaccc aggotggagt gcagtggtgc aatottggct cactgtagcc totgcctccg 180 aggttcaagc gattctcatg cctcagccgc cctagtagct aggattacag gcgccctcca 240 300 ccacacacag ctatttccag gtacattctt gacgctagga attcagcaaa gaataagaca 32 gttaaggtet eegatgetea taggeeteae attttagaga gggatgaatg tecaataage 360 33 atataaacat ataatatgtc agggtcgtat gactacaagg aacagtgatt gttacaaccc 420 34 agatgagagg gaaaaataaa ggattccaaa tatccccctt gggaagtaga gtcaggattc 480 35 aaacaaagaa ctgtatggct tcaagttcat ggtctttaat ctcctggagg ctgtctctct 540 ttctttttc ttttttaa tcagtgttgg gatcaaattc tggctcccct aggaagcatc 600 660 tggcaaggtt tcgggagcca tcgggttggc catgttatgc tggaatattt ataagcaccg 37 38 gagggttatc cccatgtcgt agaaaatgaa actgaagctc agagagattt gcactctctg 720 cccttttgta caactcattt ttccccagta tgtggaattg agggagcttc acgcttctag 780 840 40 ctgtcatgat tccaagattc tacgacatgt gggagaggat cctaaggttc ggggaaccgc 900 ggaggtttcg gggttctaga aatccgaggt tctaagccta ggtgctccaa taaacccagt gagagecage ecaggtttee ggtetgtace egetggtgea ageceagaga caageaggeg 960 43 ccaeccatga gcccctctgc ggccccctcc cgggtcccac ctcgcaggcc agctggaggg 1020 1080 egegateetg gegteeeeg aeggeetggg geeeeaatee agaggeetgg gtgggagggg 1140 45 accaagggtg tagtaaggaa gcgccttttg ctggagggca acggaccggg gcggggagtc 1200 46 gggagaccag agtgggagga aggcggggag tecaggttee geeceggage egaetteete 1260 ctggtcggcg gctgcagcgg ggtgagcggc ggcagcggcc ggggatcctg gagccatggg 48 gegegegege gaegeeatee tggatgeget ggagaacetg acegeegagg ageteaagaa 1320 gttcaagctg aagctgctgt cggtgccgct gcgcgagggc tacgggcgca tcccgcgggg 1380 50 cgcqctqctq tccatqqacq ccttqqacct caccgacaag ctggtcagct tctacctgga 1440 1500 gacctacggc gccgagctca ccgctaacgt gctgcgcgac atgggcctgc aggagatggc egggeagetg eaggeggeea egeaceaggg tgageegeec eegtteeeet eeaceeegte 1560 53 tttcccctcc acccacaca gegettaccc cgcgggetet tccgctttct gttcctccta 1620 1680 54 cooctaaaca aagotgotot acoggaaagg aggotococa ogottggoot acogaccaac 1740 55 gggaccccgg ccccacggcg ggaagggaag ggaaggggat cacttggccc atatccttcc 1800 56 aggetetgga geogegeeag etgggateea ggeeeeteet eagteggeag eeaageeagg 1860 57 tgaggcctcc acacccagcc cggccccacc gcactcctgc acagcctgca tctgtgctcc

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/691,763B

DATE: 04/17/2001
TIME: 13:21:40

Input Set : A:\E03557003.txt
Output Set: N:\CRF3\04172001\1691763B.raw

·	
60 gtectggtgg cegeceetg gageeetgee eetaggettg cagaggaatt eetgaagaae	2040
61 tcaagttcag cagggacagg ccccacaccc tggctgctgg ctcatgttct cctcccaccc	2100
62 ccaggcctgc actttataga ccagcaccgg gctgcgctta tcgcgagggt cacaaacgtt	2160
63 gagtggetge tggatgetet gtaegggaag gteetgaegg atgageagta ceaggeagtg	2220
64 cgggccgagc ccaccaaccc aagcaagatg cggaagetet tcagttteac accagcetgg	2280
65 aactggacct gcaaggactt gctcctccag gccctaaggg agtcccagtc ctacctggtg	2340
66 gaggacetgg ageggagetg aggeteette ceageaacae teeggteage eeetggeaat	2400
67 cccaccaaat catcctgaat ctgatctttt tatacacaat atacgaaaag ccagcttgaa	2460
68 cttgtgtgtt ttcctgcttc tagcctgctg gcatgtgcag agctcagcta tgcttcagag	2520
69 gccacccage etecagetee atgteectag ggtetetgge acceeaaatg etteceecat	2580
70 ccttcctggt atcgccatgg aatatccctc ctcattcacc aggtggtgct cctccagtgc	2640
71 tccctaaagg gtctaacctt accattatag ataacagcct gtgacccagg tccgaaggtt	2700
72 aaaaqaqqca tqtaccaaaq qqcqcaaact qqtgggcagc tctgtccaag ccatttagaa	2760
73 acacactagt cttcatagct cccctacctt ccacattttc cactggaaga aaaaatggca	2820
74 a	2821
76 <210> SEQ ID NO: 2	
77 <211> LENGTH: 770	
78 <212> TYPE: DNA	
79 <213> ORGANISM: Homo Sapiens	
81 <220> FEATURE:	
82 <221> NAME/KEY: CDS	
83 <222> LOCATION: (75)(662)	
85 <400> SEQUENCE: 2	
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87 gggatcctgg agec atg ggg cgc gcg cgc gac gcc atc ctg gat gcg ctg	110
88 Met Gly Arg Ala Arg Asp Ala Ile Leu Asp Ala Leu	
89 1 5 10	
91 gag aac ctg acc gcc gag gag ctc aag aag ttc aag ctg aag ctg ctg	158
92 Glu Asn Leu Thr Ala Glu Glu Leu Lys Lys Phe Lys Leu Lys Leu Leu	200
93 15 20 25	
95 tog gtg cog ctg cgc gag ggc tac ggg cgc atc ccg cgg ggc gcg ctg	206
96 Ser Val Pro Leu Arg Glu Gly Tyr Gly Arg Ile Pro Arg Gly Ala Leu	200
97 30 35 40	
99 ctg tcc atg gac gcc ttg gac ctc acc gac aag ctg gtc agc ttc tac	254
100 Leu Ser Met Asp Ala Leu Asp Leu Thr Asp Lys Leu Val Ser Phe Tyr	231
100 Lett Sel Met Asp Ala Lett Asp Lett III Asp Lys Lett Val Sel IIIc 1/1 101 45 50 55 60	
101 43	302
103 ctg gag acc tac ggc gcc gag ctc acc gct aac gtg ctg cgc gac atg	302
104 Leu Glu Thr Tyr Gly Ala Glu Leu Thr Ala Asn Val Leu Arg Asp Met 105 70 75	
103	350
107 ggc ctg cag gag atg gcc ggg cag ctg cag gcg gcc acg cac cag ggc	330
108 Gly Leu Gln Glu Met Ala Gly Gln Leu Gln Ala Ala Thr His Gln Gly	
103	398
111 tet gga gee geg eea get ggg ate eag gee eet eet eag teg gea gee	390
112 Ser Gly Ala Ala Pro Ala Gly Ile Gln Ala Pro Pro Gln Ser Ala Ala	
113 95 100 105	110
115 aag cca ggc etg cac ttt ata gac cag cac egg get geg ett ate geg	446
116 Lys Pro Gly Leu His Phe Ile Asp Gln His Arg Ala Ala Leu Ile Ala	
117 110 115 120	404
119 agg gtc aca aac gtt gag tgg ctg ctg gat gct ctg tac ggg aag gtc	494

RAW SEQUENCE LISTING DATE: 04/17/2001 PATENT APPLICATION: US/09/691,763B TIME: 13:21:40

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120	_	Val	Thr	Asn	Val	_	Trp	Leu	Leu	Asp	Ala 135	Leu	Tyr	Gly	Lys	Val 140	
121	125					130						~~~	000	200	330		542
123	ctg	acg	gat	gag	cag	tac	cag	gca	gra	cgg	315	Clu	Dro	Thr	λan	Dro	342
124	Leu	Thr	Asp	GLU		Tyr	GIn	Ala	Val		нта	GIU	PIO	1111		FIO	
125					145					150			٠		155	200	590
127	agc	aag	atg	cgg	aag	ctc	ttc	agt	ttc	aca	cca	gee	Egg	aac	Lyy	mbm	330
128	Ser	Lys	Met		Lys	Leu	Phe	Ser		Inr	Pro	Ата	Trp		тгр	THE	
129				160					165					170			620
131	tgc	aag	gac	ttg	ctc	ctc	cag	gcc	cta	agg	gag	tcc	cag	tcc	tac	ctg	638
132	Cys	Lys	_	Leu	Leu	Leu	Gln			Arg	GIu	Ser		ser	туг	Leu	
133	175 180/ 185 gtg gag gac ctg gag cgg agc tga ggctccttcc cagcaacact ccggtcagcc										600						
135									ggct	tcct	tee (cage	aaca	ct co	eggto	cagcc	692
136	Val	Glu	Asp	Leu	Glu	Arg		*									
137		190					195										750
139	cctggcaatc ccaccaaatc atcctgaatc tgatcttttt atacacaata tacgaaaagc												752				
140													770				
142	<210	> SE	Q ID	NO:	3												
143	<211	> LE	NGTH	: 19	5												
144	<212	> TYl	PE: 1	PRT													
145	<213	> OR	GANIS	SM: 1	Omo	Sap	iens										
147	<400																
148	Met	Gly	Arg	Ala	Arg	Asp	Ala	Ile	Leu	Asp	Ala	Leu	Glu	Asn	Leu	Thr	
149	1				5					10					15		
150	Ala	Glu	Glu	Leu	Lys	Lys	Phe	Lys	Leu	Lys	Leu	Leu	Ser	Val	Pro	Leu	
151				20					25					30			
152	Arq	Glu	Gly	Tyr	Gly	Arg	Ile	Pro	Arg	Gly	Ala	Leu	Leu	Ser	Met	Asp	
153			35	_	_			40					45				
154	Ala	Leu	Asp	Leu	Thr	Asp	Lys	Leu	Val	Ser	Phe	Tyr	Leu	Glu	Thr	Tyr	
155		50	_				55					60					
156	Gly	Ala	Glu	Leu	Thr	Ala	Asn	Val	Leu	Arg	Asp	Met	Gly	Leu	Gln	Glu	
157	65					70					75					80	
158	Met	Ala	Gly	Gln	Leu	Gln	Ala	Ala	Thr	His	Gln	Gly	Ser	Gly	Ala	Ala	
159			•		85					90					95		
160	Pro	Ala	Glv	Ile	Gln	Ala	Pro	Pro	Gln	Ser	Ala	Ala	Lys	Pro	Gly	Leu	
161			•	100					105					110			
162	His	Phe	Ile	Asp	Gln	His	Arq	Ala	Ala	Leu	Ile	Ala	Arg	Val	Thr	Asn	
163			115	•				120					125				
164	Val	Glu	Trp	Leu	Leu	Asp	Ala	Leu	Tyr	Gly	Lys	Val	Leu	Thr	Asp	Glu	
165		130				1	135		-	-	•	140					
166	Gln	Tyr	Gln	Ala	Val	Arq	Ala	Glu	Pro	Thr	Asn	Pro	Ser	Lys	Met	Arg	
167	145	-1-	0	••-		150					155					160	
168		Leu	Phe	Ser	Phe	Thr	Pro	Ala	Trp	Asn	Trp	Thr	Cys	Lys	Asp	Leu	
169	-10				165				•	170	-		-	-	175		
170	Len	Leu	Gln	Ala		Ara	Glu	Ser	Gln		Tyr	Leu	Val	Glu	Asp	Leu	
171	_cu			180		3			185		4 .			190	-		
172	Glu	Arg	Ser	200													
173	O_Lu	y	195														
	175 <210> SEQ ID NO: 4																
176 <211> LENGTH: 626																	
1,0	~211.	التالب -	.,0111	. 02	•												

RAW SEQUENCE LISTING DATE: 04/17/2001 PATENT APPLICATION: US/09/691,763B TIME: 13:21:40

Input Set : A:\E03557003.txt

Output Set: N:\CRF3\04172001\1691763B.raw

```
177 <212> TYPE: DNA
178 <213> ORGANISM: Homo Sapiens
180 <400> SEQUENCE: 4
181 agegeetttt getggaggge aacggacegg ggeggggagt egggagaeca gagtgggagg
                                                                                                                                                120
182 aaggegggga gtecaggtte egeeeeggag eegactteet eetggtegge ggetgeageg
183 gggtgagegg eggeagegge eggggateet ggageeatgg ggegegege egaegeeate
184 ctggatgcgc tggagaacct gaccgccgag gagctcaaga agttcaagct gaagctgctg
                                                                                                                                                240
185 teggtgeege tgegegaggg etaegggege atecegeggg gegegetget gteeatggae
                                                                                                                                                300
186 gccttggacc tcaccgacaa gctggtcagc ttctacctgg agacctacgg cgccgagctc
                                                                                                                                                360
187 acceptaace tecteced categories categorie
                                                                                                                                                420
                                                                                                                                                480
189 agegettace eegeggete tteegettte tgtteeteet acceetaaac aaagetgete
190 taccggaaag gaggctcccc acgcttggcc taccgaccaa cgggaccccg gcccacggc
                                                                                                                                                600
191 gggaagggaa gggaagggga tcactt
                                                                                                                                                626
193 <210> SEQ ID NO: 5
194 <211> LENGTH: 340
195 <212> TYPE: DNA
196 <213> ORGANISM: Homo Sapiens
198 <220> FEATURE:
199 <221> NAME/KEY: CDS
200 <222> LOCATION: (67)...(339)
202 <400> SEQUENCE: 5
                                                                                                                                                60
203 ccgacticct cctggtcggc ggctgcagcg gggtgagcgg cggcagcggc cggggatcct
204 ggagec'atg ggg ege gee gae gee ate etg gat geg etg gag aac
                                                                                                                                              108
205
                      Met Gly Arg Ala Arg Asp Ala Ile Leu Asp Ala Leu Glu Asn
206
                       1
                                                      5
                                                                                           10
208
         ctg acc gcc gag gag ctc aag aag ttc aag ctg aag ctg tcg gtg
                                                                                                                                              156
209 Leu Thr Ala Glu Glu Leu Lys Lys Phe Lys Leu Lys Leu Ser Val
210
                                                 20
                                                                                                                                              204
212 ccg ctg cgc gag ggc tac ggg cgc atc ccg cgg ggc gcg ctg ctg tcc
213 Pro Leu Arg Glu Gly Tyr Gly Arg Ile Pro Arg Gly Ala Leu Leu Ser
214
                                                                                40
216 atg gac gcc ttg gac ctc acc gac aag ctg gtc agc ttc tac ctg gag
                                                                                                                                              252
217 Met Asp Ala Leu Asp Leu Thr Asp Lys Leu Val Ser Phe Tyr Leu Glu
218
                                 50
                                                                        55
220 acc tac ggc gcc gag ctc acc gct aac gtg ctg cgc gac atg ggc ctg
                                                                                                                                              300
221
        Thr Tyr Gly Ala Glu Leu Thr Ala Asn Val Leu Arg Asp Met Gly Leu
222
                          65
                                                               70
224 cag gag atg gcc ggg cag ctg cag gcg gcc acg cac cag g
                                                                                                                                              340
225 Gln Glu Met Ala Gly Gln Leu Gln Ala Ala Thr His Gln
226
                 80
                                                        85
229 <210> SEQ ID NO: 6
230 <211> LENGTH: 91
231 <212> TYPE: PRT
232 <213> ORGANISM: Homo Sapiens
234 <400> SEQUENCE: 6
235 Met Gly Arg Ala Arg Asp Ala Ile Leu Asp Ala Leu Glu Asn Leu Thr
                                                                             10
237 Ala Glu Glu Leu Lys Lys Phe Lys Leu Lys Leu Leu Ser Val Pro Leu
```

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Output Set: N:\CRF3\04172001\1691763B.raw

```
20
238
239 Arg Glu Gly Tyr Gly Arg Ile Pro Arg Gly Ala Leu Leu Ser Met Asp
             35
                                 40
                                                      45
241
     Ala Leu Asp Leu Thr Asp Lys Leu Val Ser Phe Tyr Leu Glu Thr Tyr
         50
                             55
                                                  60
242
243
     Gly Ala Glu Leu Thr Ala Asn Val Leu Arg Asp Met Gly Leu Gln Glu
                         70
244
    65
                                              75
245 Met Ala Gly Gln Leu Gln Ala Ala Thr His Gln
                     85
248 <210> SEQ ID NO: 7
249 <211> LENGTH: 57
250 <212> TYPE: DNA
251 <213> ORGANISM: Homo Sapiens
253 <220> FEATURE:
254 <221> NAME/KEY: CDS
255 <222> LOCATION: (3)...(56)
257 <400> SEQUENCE: 7
258 gc tet gga gcc gcg cca gct ggg atc cag gcc cct cct cag tcg gca
                                                                            47
        Ser Gly Ala Ala Pro Ala Gly Ile Gln Ala Pro Pro Gln Ser Ala
259
262 gcc aag cca' g
                                                                            57
263 Ala Lys Pro
267 <210> SEQ ID NO: 8
268 <211> LENGTH: 18
269 <212> TYPE: PRT
270 <213> ORGANISM: Homo Sapiens
272 <400> SEQUENCE: 8
273 Ser Gly Ala Ala Pro Ala Gly Ile Gln Ala Pro Pro Gln Ser Ala Ala
274
                                         10
275 Lys Pro
278 <210> SEQ ID NO: 9
279 <211> LENGTH: 356
280 <212> TYPE: DNA
281 <213> ORGANISM: Homo Sapiens
283 <220> FEATURE:
284 <221> NAME/KEY: CDS
285 <222> LOCATION: (3)...(258)
287 <400> SEQUENCE: 9
     gc ctg cac ttt ata gac cag cac cgg gct gcg ctt atc gcg agg gtc
                                                                            47
289
        Leu His Phe Ile Asp Gln His Arg Ala Ala Leu Ile Ala Arg Val
                        5
290
        1
                                             10
                                                                  15
292 aca aac gtt gag tgg ctg ctg gat gct ctg tac ggg aag gtc ctg acg
                                                                            95
293 Thr Asn Val Glu Trp Leu Leu Asp Ala Leu Tyr Gly Lys Val Leu Thr
294
                      20
                                          25
                                                              30
296
     gat gag cag tac cag gca gtg cgg gcc gag ccc acc aac cca agc aag
                                                                           143
297 Asp Glu Gln Tyr Gln Ala Val Arg Ala Glu Pro Thr Asn Pro Ser Lys
298
                                      40
                                                          45
300 atg cgg aag etc ttc agt ttc aca eca gec tgg aac tgg ace tgc aag
301 Met Arg Lys Leu Phe Ser Phe Thr Pro Ala Trp Asn Trp Thr Cys Lys
```



Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/691,763B TIME: 13:21:41

DATE: 04/17/2001

Input Set : A:\E03557003.txt

Output Set: N:\CRF3\04172001\1691763B.raw

 $L\!:\!537$ $M\!:\!341$ W: (46) "n" or "Xaa" used, for SEQ ID#:22

L:581 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:23

L:590 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23